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By: Nancy Ramos Printed: Nancy Ramos

# IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Bandman et al.

Title: HUMAN EXTRACELLULAR MATRIX PROTEINS

Serial No.: To Be Assigned Filing Date: Herewith

Examiner: To Be Assigned Group Art Unit: To Be Assigned

**Official Draftsman**

Commissioner for Patents

Washington, D.C. 20231

## SUBMISSION OF FORMAL DRAWINGS

Sir:

Transmitted herewith are Figure(s) 1A, 1B, 1C, 1D, 1E, 1F, 1G, 2A, 2B, 2C, 2D, 2E, 3A, 3B, 4A, 4B, and 4C, as seventeen (17) sheets of formal drawings for this application. Each sheet of drawing indicates the identifying indicia suggested in 37 CFR Section 1.84(c) on the reverse side of the drawings.

Applicants believe that no fee is due with this paper. However, if the Commissioner determines that a fee is necessary, the Commissioner is hereby authorized to charge any additional fees associated with this communication or credit any overpayment to Deposit Account No. **09-0108**. **A duplicate copy of this communication is enclosed.**

If there are any questions regarding the above, the Examiner is invited to call the undersigned at 650-855-0555.

Respectfully submitted,

INCYTE GENOMICS, INC.

Date: 16<sup>th</sup> April 2001

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CCA	AGA	TTG	TTG	TGA	GTC	TAG	CCA	GTT	GAG	CGC	TGT	AAT	CTG	AAC	CAG	54
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																954
																963
																972
																981
																990
																1000

FIGURE 1A



819	TGC ATC AAT ACT GAA GGC GGC TAC ACC TGC TCC TGC ACC GAC GAT TAT TGG CTT	837	846	855	864
C I N T E G G Y T C S C T D G Y W L					
873	CTG GAA GGC CAG TGC TTA GAC ATT GAT GAA TGT CGC TAT GGT TAC TGC CAG CAG	891	900	909	918
L E G Q C L D I D E C R Y G Y C Q Q					
927	CTC TGT GCG AAT GTT CCT GGA TCC TAT TCT TGT ACA TGC AAC CCT GGT TTT ACC	945	954	963	972
L C A N V P G S Y S C T C N P G F T					
981	CTC AAT GAG GAT GGA AGG TCT TGC CAA GAT GTG AAC GAG TGT GCC ACC GAG AAC	999	1008	1017	1026
L N E D G R S C Q D V N E C A T E N					
1035	CCC TGC GTG CAA ACC TGC GTC AAC ACC TAC GGC TCT TTC ATC TGC CGC TGT GAC	1053	1062	1071	1080
P C V Q T C V N T Y G S F I C R C D					
1089	CCA GGA TAT GAA CTT GAG GAA GAT GGC GTT CAT TGC AGT GAT ATG GAC GAG TGC	1107	1116	1125	1134
P G Y E L E E D G V H C S D M D E C					
1143	AGC TTC TCT GAG TTC CTC TGC CAA CAT GAG TGT GTG AAC CAG CCC GGC ACA TAC	1161	1170	1179	1188
S F S E F L C Q H E C V N Q P G T Y					

FIGURE 1C

1197	1206	1215	1224	1233	1242
TTC TGC TCC	TGC CCT CCA	TAC ATC CTG	CTG GAT GAC	AAC CGA AGC	TGC CAA
F C S C	P P G Y	I L L D	D N R S	C C Q	
1251	1260	1269	1278	1287	1296
GAC ATC AAC	GAA TGT GAG	CAC AGG AAC	CAC ACG TGC	AAC CTG CAG	CAG ACG TGC
D I N E	C E H R	N H T C	N L Q	Q T C	
1305	1314	1323	1332	1341	1350
TAC AAT TTA	CAA GGG GGC	TTC AAA TGC	ATC GAC CCC	ATC CGC TGT	GAG GAG CCT
Y N L Q	G G F K	C I D P	I R C E	E P	
1359	1368	1377	1386	1395	1404
TAT CTG AGG	ATC AGT GAT	AAC CGC TGT	ATG TGT CCT	GCT GAG AAC	CCT GGC TGC
Y L R I	S D N R	C M C P	A E N P	G C	
1413	1422	1431	1440	1449	1458
AGA GAC CAG	CCC TTT ACC	ATC TTG TAC	CGG GAC ATG	GAC GTG GTG	TCA GGA CGC
R D Q P	F T I L	Y R D M	D V V S	G R	
1467	1476	1485	1494	1503	1512
TCC GTT CCC	GCT GAC ATC	TTC CAA ATG	CAA GCC ACG	ACC CGC TAC	CCT GGG GCC
S V P A	D I F Q	M Q A T	R Y P G	A	
1521	1530	1539	1548	1557	1566
TAT TAC ATT	TTT CAG ATC	AAA TCT GGG	AAT GAG GGC	AGA GAA TTT	TAC ATG CGG
Y Y I F	Q I K S	G N E G	R E F Y	M R	

FIGURE 1D

1575	1584	1593	1602	1611	1620
CAA ACG GGC CCC ATC AGT GCC ACC CTG GTG ATG ACA CGC CCC ATC AAA GGG CCC					
Q T G P I S A T L V M T R P I K G P					
1629	1638	1647	1656	1665	1674
CGG GAA ATC CAG CTG GAC TTG GAA ATG ATC ACT GTC AAC ACT GTC ATC AAC TTC					
R E I Q L D L L E M I T V N T V I N F					
1683	1692	1701	1710	1719	1728
AGA GGC AGC TCC GTG ATC CGA CTG CGG ATA TAT GTG TCG CAG TAC CCA TTC TGA					
R G S S V I R L R I Y V S Q Y P F					
1737	1746	1755	1764	1773	1782
GCC TCG GGC TGG AGC CTC CGA CGC TGC CTC TCA TTG GCA CCA AGG GAC AGG AGA					
AGA GAG GAA ATA ACA GAG AGA ATG AGA GCG ACA CAG ACG TTA GGC ATT TCC TGC					
1791	1800	1809	1818	1827	1836
TGA ACG TTT CCC CGA AGA GTC AGC CCC GAC TTC CTG ACT CTC ACC TGT ACT ATT					
1845	1854	1863	1872	1881	1890
GCA GAC CTG TCA CCC TGC AGG ACT TGC CAC CCC CAG TTC CTA TGA CAC AGT TAT					
1899	1908	1917	1926	1935	1944
CAA AAA GTA TTA TCA TTG CTC CCC TGA TAG AAG ATT GTT GGT GAA TTT TCA AGG					
1953	1962	1971	1980	1989	1998

FIGURE 1E

2007	2016	2025	2034	2043	2052
CCT TCA GTT TAT TTC CAC TAT TTT CAA AGA AAA TAG ATT AGG TTT GCG GGG GTC					
2061	2070	2079	2088	2097	2106
TGA GTC TAT GTT CAA AGA CTG TGA ACA GCT TGC TGT CAC TTC TTC ACC TCT TCC					
2115	2124	2133	2142	2151	2160
ACT CCT TCT CTC CTC ACT GTG TTA CTG CTT TGC AAA GAC CCG GGA GCT GGC GGG GAA					
2169	2178	2187	2196	2205	2214
CCC TGG GAG TAG CTA GTT TGC TTT TTG CGT ACA CAG AGA AGG CTA TGT AAA CAA					
2223	2232	2241	2250	2259	2268
ACC ACA GCA GGA TCG AAG GGT TTT TAG AGA ATG TGT TTC AAA ACC ATG CCT GGT					
2277	2286	2295	2304	2313	2322
ATT TTC AAC CAT AAA AGA AGT TTC AGT TGT CCT TAA ATT TGT ATA ACG GTT TAA					
2331	2340	2349	2358	2367	2376
TTC TGT CTT GTT CAT TTT GAG TAT TTT TAA AAA ATA TGT CGT AGA ATT CCT TCG					
2385	2394	2403	2412	2421	2430
AAA GGC CTT CAG ACA CAT GCT ATG TTC TGT CTT CCC AAA CCC AGT CTC CTC TCC					
2439	2448	2457	2466	2475	2484
ATT TTA GCC CAG TGT TTT CTT TGA GGA CCC CTT AAT CTT GCT TTC TTT AGA ATT					

FIGURE 1F

TTT ACC CAA TTG GAT TGG AAT GCA GAG 2511 2520 2529 2538  
AGA AAA AAA AAA

FIGURE 1G



9	18	27	36	45	54
TG GGT GCA AGC TCA CAA CCG TAA CAG CCA CCA AAG CTTT CAG TGG CCG GCC					
63	72	81	90	99	108
CTT CAC ATC CAG ACT TGC CTG AGA GGA CCC ACC TCT GAG TGT CCA GTG GTC AGT					
117	126	135	144	153	162
TGC CCC AGG ATG GGG ACC ACA GCC AGA GCA GCC TTG GTC TTG ACC TAT TTG GCT					
M G T T A R A A L V L T Y L A					
171	180	189	198	207	216
GTT GCT TCT GCT GCC TCT GAG GGA GGC TTC ACG GCT ACA GGA CAG AGG CAG CTG					
V A S A A S E G G F T A T G Q R Q L					
225	234	243	252	261	270
AGG CCA GAG CAC TTT CAA GAA GTT GGC TAC GCA GCT CCC CCC TCC CCA CCC CTA					
R P E H F Q E V G Y A A P P S P L					
279	288	297	306	315	324
TCC CGA AGC CTC CCC ATG GAT CAC CCT GAC TCC TCT CAG CAT GGC CCT CCC TTTT					
S R S L P M D H P D S S Q H G P P F					
333	342	351	360	369	378
GAG GGA CAG AGT CAA GTG CAG CCC CCT TCT CAG GAG GCC ACC CCT CTC CAA					
E G Q S Q V Q P P P S Q E A T P L Q					

FIGURE 2A

387	396	405	414	423	432
CAG GAA AAG CTG CTA CCT GCC CAA CTC CCT GCT GAA AAG GAA GTG GGT CCC CCT					
Q E K L L P A Q L P A E K E V G P P					
441	450	459	468	477	486
CTC CCT CAG GAA GCT GTC CCC CTC CAA AAA GAG CTG CCC TCT CTC CAG CAC CCC					
L P Q E A V P L Q K E L P S L Q H P					
495	504	513	522	531	540
AAT GAA CAG AAG GAA GGA ATG CCA GCT CCA TTT GGG GAC CAG AGC CAT CCA GAA					
N E Q K E G M P A P F G D Q S H P E					
549	558	567	576	585	594
CCT GAG TCC TGG AAT GCA GCC CAG CAC TGC CAA CAG GAC CGG TCC CAA GGG GGC					
P E S W N A A Q H C Q Q D R S Q G G					
603	612	621	630	639	648
TGG GGC CAC CGG CTG GAT GGC TTC CCC CCT GGG CGG CCT TCT CCA GAC AAT CTG					
W G H R L D G F P P G R P S P D N L					
657	666	675	684	693	702
AAC CAA ATC TGC CTT CCT AAC CGT CAG CAT GTG GTA TAT GGT CCC TGG AAC CTA					
N Q I C L P N R Q H V V Y G P W N L					
711	720	729	738	747	756
CCA CAG TCC AGC TAC TCC CAC CTC ACT CGC CAG GGT GAG ACC CTC AAT TTC CTG					
P Q S S Y S H L T R Q G E T L N F L					

FIGURE 2B

765	774	783	792	801	810
GAG ATT GGA TAT TCC CGC TGC TGC CAC TGC CGC AGC CAC ACA AAC CGC CTA GAG					
E I G Y S R C C H C R S H T N R L E					
819	828	837	846	855	864
TGT GCC AAA CTT GTG TGG GAG GAA GCA ATG AGC CGA TTC TGT GAG GCC GAG TTC					
C A K L V W E E A M S R F C E A E F					
873	882	891	900	909	918
TCG GTC AAG ACC CGA CCC CAC TGG TGC TGC ACG CGG CAG GGG GAG GCT CGG TTC					
S V K T R P H W C C T R Q G E A R F					
927	936	945	954	963	972
TCC TGC TTC CAG GAG GAA GCT CCC CAG CCA CAC TAC CAG CTC CGG GCC TGC CCC					
S C F Q E E A P Q P H Y Q L R A C P					
981	990	999	1008	1017	1026
AGC CAT CAG CCT GAT ATT TCC TCG GGT CTT GAG CTG CCT TTC CCT CCT GGG GTG					
S H Q P D I S S G L E L P F P P G V					
1035	1044	1053	1062	1071	1080
CCC ACA TTG GAC AAT ATC AAG AAC ATC TGC CAC CTG AGG CGC TTT CGC TCT GTG					
P T L D N I K N I C H L R R F R S V					
1089	1098	1107	1116	1125	1134
CCA CGC AAC CTG CCA GCT ACT GAC CCC CTA CAA AGG GAG CTG CTG GCA CTG ATC					
P R N L P A T D P L Q R E L L A L I					

FIGURE 2C

1143	1152	1161	1170	1179	1188
CAG CTG GAG AGG GAG TTC CAG CGC TGC TGC CGC CAG GGG AAC AAT CAC ACC TGT					
Q L E R E F Q R C C R Q G N N H T C					
1197	1206	1215	1224	1233	1242
ACA TGG AAG GCC TGG GAG GAT ACC CTT GAC AAA TAC TGT GAC CGG GAG TAT GCT					
T W K A W E D T L D K Y C D R E Y A					
1251	1260	1269	1278	1287	1296
GTG AAG ACC CAC CAC CAC TTG TGT TGC CGC CAC CCT CCC AGC CCT ACT CGG GAT					
V K T H H H L C C R H P P S P T R D					
1305	1314	1323	1332	1341	1350
GAG TGC TTT GCC CGT CGG GCT CCT TAC CCC AAC TAT GAC CGG GAC ATC TTG ACC					
E C F A R R A P Y P N Y D R D I L T					
1359	1368	1377	1386	1395	1404
ATT GAC ATC GGT CGA GTC ACC CCC AAC CTC ATG GGC CAC CTC TGT TGA AAC CAA					
I D I G R V T P N L M G H L C G N Q					
1413	1422	1431	1440	1449	1458
AGA GTT CTC ACC AAG CAT AAA CAT ATT CCT GGG CTG ATC CAC AAC ATG ACT GCC					
R V L T K H K H I P G L I H N M T A					
1467	1476	1485	1494	1503	1512
CGC TGC TGT GAC CTG CCA TTT CCA GAA CAG GCC TGC TGT TGA GAG GAG AAA					
R C C D L P P F P E Q A C C A E E E K					

FIGURE 2D

1521	1530	1539	1548	1557	1566
TTA ACC TTC ATC AAT GAT CTG TGT GGT CCC CGA CGT AAC ATC TGG CGA GAC CCT					
L T F I N D L C G P R R N I W R D P					
1575	1584	1593	1602	1611	1620
GCC CTC TGC TGT TAC CTG AGT CCT GGG GAT GAA CAG GTC AAC TGC TTC AAC ATC					
A L C C Y L S P G D E Q V N C F N I					
1629	1638	1647	1656	1665	1674
AAT TAT CTG AGG AAC GTG GCT CTA GTG TCT GGA GAC ACT GAG AAC GCC AAG GGC					
N Y L R N V A L V S G D T E N A K G					
1683	1692	1701	1710	1719	1728
CAG GGG GAG CAG GGC TCA ACT GGA GGA ACA AAT ATC AGC TCC ACC TCT GAG CCC					
Q G E Q G S T G G T N I S S T S E P					
1737	1746	1755	1764	1773	1782
AAG GAA GAA TGA GTC ACC CCA GAG CCC TAG AGG GTC AGA TGG GGG GAA CCC CAC					
K E E					
1791	1800	1809	1818	1827	1836
CCT GCC CCA CCC ATC TGA ACA CTC ATT ACA CTA AAC ACC TCT TGG ATT TGG TGT					
1845	1854	1863	1872	1881	1890
CCT CAT TGT CTA TCT AAT GTC TCA CCC GCA GTG TTT TAA GTG GAT CTT GGT GCC					
1899					
CTG GCC CAG G					

FIGURE 2E

1 M P G I K R I L T V T I L A L C L P S P 45517  
1 M A T S G V L P G G G F V A S A A A V A GI 458228

21 G N A Q A Q C T N G F D L D R Q S G Q C 45517  
21 G P E M Q T G R N N F V I R R N P A D - GI 458228

41 L D I D E C R T I P E A C R G D M M C V 45517  
40 - - - - - - - - - - - - - - - - GI 458228

61 N Q N G G Y L C I P R T N P V Y R G P Y 45517  
40 - - - - - - - - - - - - - - - - GI 458228

81 S N P Y S T P Y S G P Y P A A A P P L S 45517  
40 - - - - - - - - - - - - - - - P Q R GI 458228

101 A P N Y P T I S R P L I C R F G Y Q M D 45517  
43 I P S N P - - S H R I Q C A A G Y E Q S GI 458228

121 E S N Q C V D V D E C A T D S H Q C N P 45517  
61 E H N V C Q D I D E C T A G T H N C R A GI 458228

141 T Q I C I N T E G G Y T C S C T D G Y W 45517  
81 D Q V C I N L R G S F A C Q C P P G Y Q GI 458228

161 L L E G Q C L D I D E C R Y G - Y C Q Q 45517  
101 K R G E Q C V D I D E C T I P P Y C H Q GI 458228

180 L C A N V P G S Y S C T C N P G F T L N 45517  
121 R C V N T P G S F Y C Q C S P G F Q L A GI 458228

200 E D G R S C Q D V N E C A T E N P C V Q 45517  
141 A N N Y T C V D I N E C D A S N Q C A Q GI 458228

220 T C V N T Y G S F I C R C D P G Y E L E 45517  
161 Q C Y N I L G S F I C Q C N Q G Y E L S GI 458228

FIGURE 3A

240 E D G V H C S D M D E C S F S E F L C Q 45517  
181 S D R L N C E D I D E C R T S S Y L C Q GI 458228

260 H E C V N Q P G T Y F C S C P P G Y I L 45517  
201 Y Q C V N E P G K F S C M C P Q G Y Q V GI 458228

280 L D D N R S C Q D I N E C E H R N H T C 45517  
221 V R - S R T C Q D I N E C E T T N E - C GI 458228

300 N L Q Q T C Y N L Q G G F K C I D P I R 45517  
239 R E D E M C W N Y H G G F R C Y P R N P GI 458228

320 C E E P Y L R I S D N R C M C P A E N P 45517  
259 C Q D P Y I L T P E N R C V C P V S N A GI 458228

340 G C R D Q P F T I L Y R D M D V V S G R 45517  
279 M C R E L P Q S I V Y K Y M S I R S D R GI 458228

360 S V P A D I F Q M Q A T T R Y P G A Y Y 45517  
299 S V P S D I F Q I Q A T T I Y A N T I N GI 458228

380 I F Q I K S G N E G R E F Y M R Q T G P 45517  
319 T F R I K S G N E N G E F Y L R Q T S P GI 458228

400 I S A T L V M T R P I K G P R E I Q L D 45517  
339 V S A M L V L V K S L S G P R E H I V D GI 458228

420 L E M I T V N T V I N F R G S S V I R L 45517  
359 L E M L T V S S I G T F R T S S V L R L GI 458228

440 R I Y V S Q Y P F 45517  
379 T I I V G P F S F GI 458228

FIGURE 3B

1 M G T T A R A A L V L T Y L A V A S A A 1621777  
1 M G T V S R A A L I L A C L A L A S A A GI 496120

21 S E G G F T A T G Q R Q L R P E - - - 1621777  
21 S E G A F K A S D Q R E M T P E R L F Q GI 496120

37 H F Q E V G Y A A P P S P P L S R S L P 1621777  
41 H L H E V G Y A A P P S L P Q T R R L R GI 496120

57 M D H P D S S Q H G P P - F E G Q S Q V 1621777  
61 V D H S V T S L H D P P L F E E Q R E V GI 496120

76 Q P P P S Q E A T P L Q Q E K L L P A Q 1621777  
81 Q P P S S P E D I P V Y E E D W P T F L GI 496120

96 L P A E K E V G P P L P Q E A V P L Q K 1621777  
101 N P N V D K A G P A V P Q E A I P L Q K GI 496120

116 E L P S L Q - - - - - - - - - - H P 1621777  
121 E Q P P P Q V H I E Q K E I D P P A Q P GI 496120

124 N E - - - Q K E G M P A P F G D Q S H P 1621777  
141 Q E E I V Q K E V K P H T L A G Q L P P GI 496120

141 E P E S W N A A Q H C Q Q D R S Q G G W 1621777  
161 E P R T W N P A R H C Q Q G R - R G V W GI 496120

161 G H R L D G F P P G R P S P D N L N Q I 1621777  
180 G H R L D G F P P G R P S P D N L K Q I GI 496120

181 C L P N R Q H V V Y G P W N L P Q S S Y 1621777  
200 C L P E R Q H V I Y G P W N L P Q T G Y GI 496120

201 S H L T R Q G E T L N F L E I G Y S R C 1621777  
220 S H L S R Q G E T L N V L E T G Y S R C GI 496120

FIGURE 4A



221 C H C R S H T N R L E C A K L V W E E A 1621777  
240 C P C R S D T N R L D C L K L V W E D A GI 496120

241 M S R F C E A E F S V K T R P H W C C T 1621777  
260 M T Q F C E A E F S V K T R P H L C C R GI 496120

261 R Q G E A R F S C F Q E E A P Q P H Y Q 1621777  
280 L R G E E R F S C F Q K E A P R P D Y L GI 496120

281 L R A C P S H Q P D I S S G L E L P F P 1621777  
300 L R P C P V H Q N G M S S G P Q L P F P GI 496120

301 P G V P T L D N I K N I C H L R R F R S 1621777  
320 P G L P T P D N V K N I C L L R R F R A GI 496120

321 V P R N L P A T D P L Q R E L L A L I Q 1621777  
340 V P R N L P A T D A I Q R Q L Q A L T R GI 496120

341 L E R E F Q R C C R Q G N N H T C T W K 1621777  
360 L E T E F Q R C C R Q G H N N H T C T W K GI 496120

361 A W E D T L D K Y C D R E Y A V K T H H 1621777  
380 A W E G T L D G Y C E R E L A I K T H P GI 496120

381 H L C C R H P P S P T R D E C F A R R A 1621777  
400 H S C C H Y P P S P A R D E C F A H L A GI 496120

401 P Y P N Y D R D I L T I D I G R V T P N 1621777  
420 P Y P N Y D R D I L T L D L S R V T P N GI 496120

421 L M G H L C G N Q R V L T K H K H I P G 1621777  
440 L M G Q L C G S G R V L S K H K Q I P G GI 496120

441 L I H N M T A R C C D L P F P E Q A C C 1621777  
460 L I Q N M T V R C C E L P Y P E Q A C C GI 496120

FIGURE 4B

461 A E E E K L T F I N D L C G P R R N I W 1621777  
 480 G E E E K L A F I E N L C G P R R N S W GI 496120  
  
 481 R D P A L C C Y L S P G D E Q V N C F N 1621777  
 500 K D P A L C C D L S P E D K Q I N C F N GI 496120  
  
 501 I N Y L R N V A L V S G D T E N A K G Q 1621777  
 520 T N Y L R N V A L V A G D T G N A T G L GI 496120  
  
 521 G E Q G S T G G T N I S S T S E P K E E 1621777  
 540 G E Q G P T R G T D A N P A P G S K E E GI 496120

FIGURE 4C